

WHAT IS CLAIMED IS:

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1. A method for ameliorating the harmful effects of TNF in an animal, comprising administering to an animal in need of such treatment a therapeutically effective amount of a recombinant polypeptide having the ability to bind TNF, wherein said polypeptide is encoded by a nucleic acid molecule comprising a nucleotide sequence of the formula: $R^1-R^2-R^3-R^4$, wherein
- 5 R^1 is ATG, or the nucleotide sequence ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA (SEQ ID NO: 5), or is absent;
- 10 R^2 is the nucleotide sequence CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA (SEQ ID NO: 7) or is absent;
- R^3 is the nucleotide sequence of SEQ ID NO: 3; and
- R^4 is the nucleotide sequence GTT AAG GGC ACT GAG GAC TCA GGC ACC
- 15 ACA (SEQ ID NO: 9) or is absent.
2. The method of Claim 1, wherein R^1 is ATG, R^2 is the nucleotide sequence CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA (SEQ ID NO: 7), and R^4 is the nucleotide sequence GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA
- 20 (SEQ ID NO: 9).
3. The method of Claim 1, wherein R^1 is ATG, R^2 is the nucleotide sequence CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA (SEQ ID NO: 7), and R^4 is absent.
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4. The method of Claim 1, wherein R^1 is ATG, R^2 is absent, and R^4 is the nucleotide sequence GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA (SEQ ID NO: 9).
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5. The method of Claim 1, wherein R^1 is ATG, R^2 is absent, and R^4 is absent.

6. The method of Claim 1, wherein R¹ is the nucleotide sequence ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA (SEQ ID NO: 5), R² is the nucleotide sequence CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA (SEQ ID NO: 7), and R⁴ is the nucleotide sequence GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA (SEQ ID NO: 9).

7. The method of Claim 1, wherein R¹ is the nucleotide sequence ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA (SEQ ID NO: 5), R² is the nucleotide sequence CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA (SEQ ID NO: 7), and R⁴ is absent.

8. The method of Claim 1, wherein R¹ is the nucleotide sequence ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA (SEQ ID NO: 5), R² is absent, and R⁴ is the nucleotide sequence GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA (SEQ ID NO: 9).

9. The method of Claim 1, wherein R¹ is the nucleotide sequence ATG GGC CTC TCC ACC GTG CCT GAC CTG CTG CTG CCA CTG GTG CTC CTG GAG CTG TTG GTG GGA ATA TAC CCC TCA GGG GTT ATT GGA (SEQ ID NO: 5), R² is absent, and R⁴ is absent.

10. The method of Claim 1, wherein R¹ is absent, R² is the nucleotide sequence CTG GTC CCT CAC CTA GGG GAC AGG GAG AAG AGA (SEQ ID NO: 7), and R⁴ is the nucleotide sequence GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA (SEQ ID NO: 9).

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Station	Time	Temp	Wind	Pressure	Humidity	Clouds	Visibility	Remarks
101	0000	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
102	0100	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
103	0200	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
104	0300	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
105	0400	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
106	0500	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
107	0600	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
108	0700	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
109	0800	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
110	0900	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
111	1000	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
112	1100	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
113	1200	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
114	1300	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
115	1400	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
116	1500	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
117	1600	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
118	1700	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
119	1800	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
120	1900	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
121	2000	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
122	2100	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
123	2200	10.0	0.0	1013.0	75.0	0.0	10.0	Clear
124	2300	10.0	0.0	1013.0	75.0	0.0	10.0	Clear

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(SEQ ID NO: 8) or is absent;

R³ is the amino acid sequence of SEQ ID NO: 4; and

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wherein said polypeptide has:

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- a) at least one conservative amino acid substitution;
 - b) at least one amino acid substitution at a glycosylation site;
 - c) at least one amino acid substitution at a proteolytic cleavage site;
 - d) at least one amino acid substitution at a cysteine residue;
 - e) at least one amino acid deletion;
 - f) at least one amino acid insertion;
 - g) a C- and/or N-terminal truncation; or
 - h) a combination of modifications selected from the group consisting of conservative amino acid substitutions, amino acid substitutions at a glycosylation site, amino acid substitutions at a proteolytic cleavage site, amino acid substitutions at a cysteine residue, amino acid deletions, amino acid insertions, C-terminal truncation, and N-terminal truncation.
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16. The method of Claim 15, wherein said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$ and has at least one conservative amino acid substitution.

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17. The method of Claim 15, wherein said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$ and has at least one amino acid substitution at a glycosylation site.

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18. The method of Claim 15, wherein said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$ and has at least one amino acid substitution at a proteolytic cleavage site.

19. The method of Claim 15, wherein said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$ and has at least one amino acid substitution at a cysteine residue.

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20. The method of Claim 15, wherein said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$ and has at least one amino acid deletion.

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21. The method of Claim 15, wherein said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$ and has at least one amino acid insertion.

5 22. The method of Claim 15, wherein said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$ and has a C- and/or N-terminal truncation.

23. A method for ameliorating the harmful effects of TNF in an animal, comprising administering to an animal in need of such treatment a therapeutically effective amount of a recombinant polypeptide having the ability to bind TNF, wherein
10 said polypeptide comprises an amino acid sequence of the formula: $R^1-R^2-R^3-R^4$, wherein

R^1 is methionine, or the amino acid sequence Met Gly Leu Ser Thr Val Pro Asp Leu Leu Leu Pro Leu Val Leu Leu Glu Leu Leu Val Gly Ile Tyr Pro Ser Gly Val Ile Gly
15 (SEQ ID NO: 6), or is absent;

R^2 is the amino acid sequence Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg (SEQ ID NO: 8) or is absent;

R^3 is the amino acid sequence of SEQ ID NO: 4; and

R^4 is the amino acid sequence Val Lys Gly Thr Glu Asp Ser Gly Thr Thr (SEQ ID NO: 10) or is absent.
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24. The method of Claim 23, wherein R^1 is methionine, R^2 is the amino acid sequence Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg (SEQ ID NO: 8), and R^4 is the amino acid sequence Val Lys Gly Thr Glu Asp Ser Gly Thr Thr (SEQ ID NO: 10).

25 25. The method of Claim 23, wherein R^1 is methionine, R^2 is the amino acid sequence Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg (SEQ ID NO: 8), and R^4 is absent.

30 26. The method of Claim 23, wherein R^1 is methionine, R^2 is absent, and R^4 is the amino acid sequence Val Lys Gly Thr Glu Asp Ser Gly Thr Thr (SEQ ID NO: 10).

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27. The method of Claim 23, wherein R¹ is methionine, R² is absent, and R⁴ is absent.

28. The method of Claim 23, wherein R¹ is the amino acid sequence Met Gly Leu Ser Thr Val Pro Asp Leu Leu Leu Pro Leu Val Leu Leu Glu Leu Leu Val Gly Ile Tyr Pro Ser Gly Val Ile Gly (SEQ ID NO: 6), R² is the amino acid sequence Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg (SEQ ID NO: 8), and R⁴ is the amino acid sequence Val Lys Gly Thr Glu Asp Ser Gly Thr Thr (SEQ ID NO: 10).

29. The method of Claim 23, wherein R¹ is the amino acid sequence Met Gly Leu Ser Thr Val Pro Asp Leu Leu Leu Pro Leu Val Leu Leu Glu Leu Leu Val Gly Ile Tyr Pro Ser Gly Val Ile Gly (SEQ ID NO: 6), R² is the amino acid sequence Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg (SEQ ID NO: 8), and R⁴ is absent.

30. The method of Claim 23, wherein R¹ is the amino acid sequence Met Gly Leu Ser Thr Val Pro Asp Leu Leu Leu Pro Leu Val Leu Leu Glu Leu Leu Val Gly Ile Tyr Pro Ser Gly Val Ile Gly (SEQ ID NO: 6), R² is absent, and R⁴ is the amino acid sequence Val Lys Gly Thr Glu Asp Ser Gly Thr Thr (SEQ ID NO: 10).

31. The method of Claim 23, wherein R¹ is the amino acid sequence Met Gly Leu Ser Thr Val Pro Asp Leu Leu Leu Pro Leu Val Leu Leu Glu Leu Leu Val Gly Ile Tyr Pro Ser Gly Val Ile Gly (SEQ ID NO: 6), R² is absent, and R⁴ is absent.

32. The method of Claim 23, wherein R¹ is absent, R² is the amino acid sequence Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg (SEQ ID NO: 8), and R⁴ is the amino acid sequence Val Lys Gly Thr Glu Asp Ser Gly Thr Thr (SEQ ID NO: 10).

33. The method of Claim 23, wherein R¹ is absent, R² is the amino acid sequence Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg (SEQ ID NO: 8), and R⁴ is absent.

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35. The method of Claim 23, wherein R¹ is absent, R² is absent, and R⁴ is absent.

37. A method for ameliorating the harmful effects of TNF in an animal, comprising administering to an animal in need of such treatment a therapeutically effective amount of a recombinant polypeptide having the ability to bind TNF, wherein said polypeptide comprises the amino acid sequence of SEQ ID NO: 4 or a C- and/or N-terminally shortened sequence thereof.

39. The method of Claim 37, wherein said polypeptide comprises a C-terminally shortened sequence of the amino acid sequence of SEQ ID NO: 4.

41. A method for ameliorating the harmful effects of TNF in an animal, comprising administering to an animal in need of such treatment a therapeutically effective amount of a recombinant polypeptide having the ability to bind TNF, wherein
30 said polypeptide consists of the amino acid sequence of SEQ ID NO: 4.

43. A method for ameliorating the harmful effects of TNF in an animal,
10 comprising administering to an animal in need of such treatment a therapeutically
effective amount of a recombinant polypeptide having the ability to bind TNF, wherein
said polypeptide consists of a C-terminally shortened sequence of the amino acid
sequence of SEQ ID NO: 4.

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25 46. The method of Claim 45, wherein said polypeptide has at least one additional amino acid at the amino-terminus.

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48. The method of Claim 45, wherein said polypeptide has at least one additional amino acid at the carboxyl-terminus.

49. A method for ameliorating the harmful effects of TNF in an animal, comprising administering to an animal in need of such treatment a therapeutically effective amount of a recombinant polypeptide having the ability to bind TNF, wherein said polypeptide is encoded by a nucleic acid which hybridizes under moderately or highly stringent conditions to the complement of the nucleic acid molecule defined in Claim 1.

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50. The method of any of Claims 1, 15, or 23, wherein said polypeptide is chemically derivatized.

51. The polypeptide of any of Claims 1, 14, 15, 23, 36, 37, 41, 42, 43, 44, or 49, wherein said polypeptide is not glycosylated.

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52. The polypeptide of any of Claims 1, 14, 15, 23, 36, 37, 41, 42, 43, 44, or 49, wherein said polypeptide is glycosylated.

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53. The polypeptide of Claim 52, wherein said polypeptide is glycosylated by a CHO cell.

54. The method of any of Claims 1, 15, or 23, wherein said recombinant polypeptide is expressed in a cultured cell *in vitro* and said recombinant polypeptide is isolated therefrom.

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55. The method of Claim 54, wherein the cultured cell is a non-human cell.

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56. The method of Claim 55, wherein the non-human cell line is a prokaryotic cell.

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61. The method of Claim 54, wherein the polypeptide is glycosylated.
62. The method of Claim 54, wherein the polypeptide is not glycosylated.